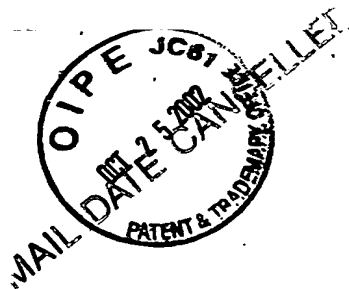


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1201 a

Figure 1



03681050.112502



MIKHFFGTVLVLLASTTIFSLDLKLIIFQQRQVNQESLKLLNKLQTL~~SI~~QQCLPH
RKNFLLPQKSLSPQQYQKGHTLAILHEMLQQIFSLFRANISLDGWEENHTEK
FLIQLHQQLEYLEALMGLEAEKLSGTLGSDNRLQVKMYFRRIHDYLENQD
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Figure 2

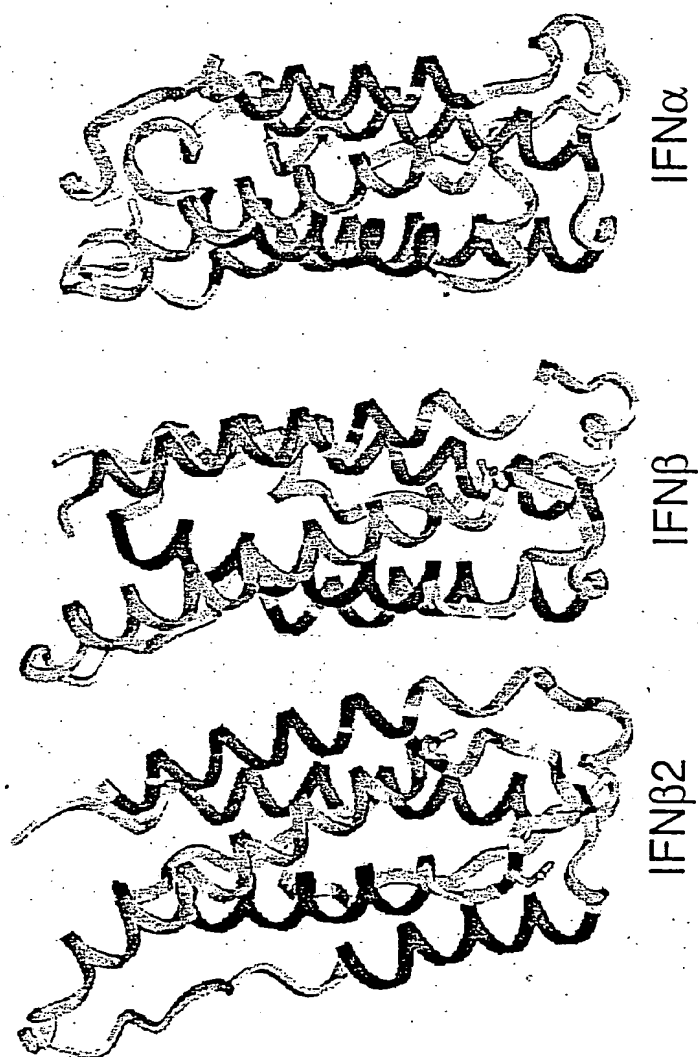


Figure 3



1

IFNB2 (1) MIEKHFFGTVLVLLASTTIFSLDLKIIIFQQRQVNOSESLLNKLQ-TESIQOCLPHRQNFLLPQKSLSP
 IFNB (1) MTNKCLEQIALLLCFSTALMSYNLLGLQRSSNFQOQKLLWQNGRLEY--CLKDRMNDIPEEIKQL
 IFNalpha8 (1) MALTFYLLVALVVLVSYSFSSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha7 (1) MARSFSLMVVLVSYSICSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha6 (1) MALPFALLMALVVLVSCKSSCSLDCDLPQTHS-LGHRRTMMLLAQMR-RISLFSCLKDRHDFEPQEEFDD
 IFNalpha5 (1) MALPFVLLMALVVLNCKSIICSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha4b (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha21 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha2 (1) MALTFALLVALVLSCKSSCSVGCGLPQTHS-LGSRRTMLLAQMR-RISLFSCLKDRHDFEPQEEFDD
 IFNalpha16 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha14 (1) MALPFALLMALVVLVSCKSSCSLGCNLSQTHS-LGNRRALILLAQMR-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha13 (1) MASPFALLMVVLVSCKSSCSLGCGLPETHS-LGNRRALILLAQMS-RISPFSCDKDRHDFEPQEEFDD
 IFNalpha10 (1) MALSFSLMAVLVLSYSICSLGCDLPQTHS-LGNRRALILLGQMG-RISPFSCDKDRHDFEPQEEFDD
 IFNomega1 (1) MALFPLLAALVMTSYSPVGSGLGCDLPQNHG-LLSRNTLVLLHQM-RISPELCLKDRHDFEPQEEFDD
 IFNgamma (1) MKYT-SYTLAFQLCIVLSGLGCYQDFYVKE----AENLKXYFNAG----H--SDVADNGTLF--LGILK
 Consensus (1) MAL F LLMALLVLS KS CSLGCDLPQTHS L NRR L LLAQM RISPFSCDKDRHDFEPQEEFDD

70

71

IFNB2 (70) QOYQKQHTLATLHEMEOQIFSLFRANISLDGWEENHTEKFLIQHQQLEYLEALMGLEAKLSGTGLSDN
 IFNB (69) QOQKEDAAALTIYEMLOIFALIFRQDSSSTGWNETIVENLLANNVYHQINHLKTVLEEKLEKEDFTRGKLM
 IFNalpha8 (69) KQFQKAQAISVLHEMIQOTFNLFSTKDSSAALDETLLEDEFYIELDQOLNDEACVIOEVGVEETPLMND
 IFNalpha7 (69) HQFQKTAQISVLHEMIQOTFNLFSTEDSSAAWEQSILKEKSTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha6 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSVANDERLLDKLYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha5 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSATWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha4b (69) HQFQKTAQISVLHEMIQOTFNLFSTEDSSAAWEQSILKEKSTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha21 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSATWEQSILKEKSTELNQQOLNDEACVIOEVGVEETPLMND
 IFNalpha2 (68) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha16 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha14 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha13 (69) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNalpha10 (69) NQFQKAQAISVLHEMIQOTFNLFSTEDSSAAWEQSILKEKSTELYQOLNDEACVIOEVGVEETPLMND
 IFNomega1 (69) SOLQKAHVMSVLHEMIQOTFNLFSTEDSSAAWDETLLDKFYTELYQOLNDEACVIOEVGVEETPLMND
 IFNgamma (58) NWKEESDRKIMQSCIVSFYFKLFKNFKD---DQS-HQKSVETIKEDMN-VKFFNSNKKKRDDFEKITYNY
 Consensus (71) NQFQKAQAISVLHEMIQOTFNLFSTKDSSAAWDE LDKF TELYQOLNDEACV QEVGVEETPLMND

140

141

IFNB2 (140) LRLOVKMYFRRIHDYLE-NQDYSTCAWATVQVEISRCFLFFVFSLTEKLSKQGRPLNDMKQELTTEFRSPR
 IFNB (139) SSLHLKRYVGRILHYLK-AKEYSHCAWTEVRVEILRNEYFINRLTGYLRLN-----
 IFNalpha8 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSLINLQKRLKSKE-----
 IFNalpha7 (139) FILAVRKYFORITLYLM-EKKYSPCAWEVRAEIMRSFSLINLQKRLRRKD-----
 IFNalpha6 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSSRNQERLRRKE-----
 IFNalpha5 (139) SILTVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSLSANLQERLRRKE-----
 IFNalpha4b (139) SILAVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSLINLQKRLRRKD-----
 IFNalpha21 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSLSKIQERLRRKE-----
 IFNalpha2 (138) SILAVRKYFORITLYLK-EKKYSPCAWEVRAEIMRSFSLSTNLQESLRSKE-----
 IFNalpha16 (139) SILAVRKYFORITLYLM-GKKYSPCAWEVRAEIMRSFSLSTNLQKGLRRKD-----
 IFNalpha14 (139) SILAVRKYFORITLYLM-EKKYSPCAWEVRAEIMRSFSLSTNLQKRLRRKD-----
 IFNalpha13 (139) SILAVRKYFORITLYLT-EKKYSPCAWEVRAEIMRSFSLSTNLQERLRRKE-----
 IFNalpha10 (139) SILAVRKYFORITLYLI-ERKYSKPCAWEVRAEIMRSFSLSTNLQKRLRRKD-----
 IFNomega1 (139) PALITRRYFQGRIVYLK-EKKYSDCAWEVRAEIMKSLFLSTNMQERLRSKDRDLGSS-----
 IFNgamma (122) SYTDLVQRKAIHELIVMAELSPAATGKR---KRSQML---FRGRASQ-----
 Consensus (141) SILAVRKYFORITLYL EKKYSPCAWEVRAEIMRSFSLSTNLQ RLRRK

210

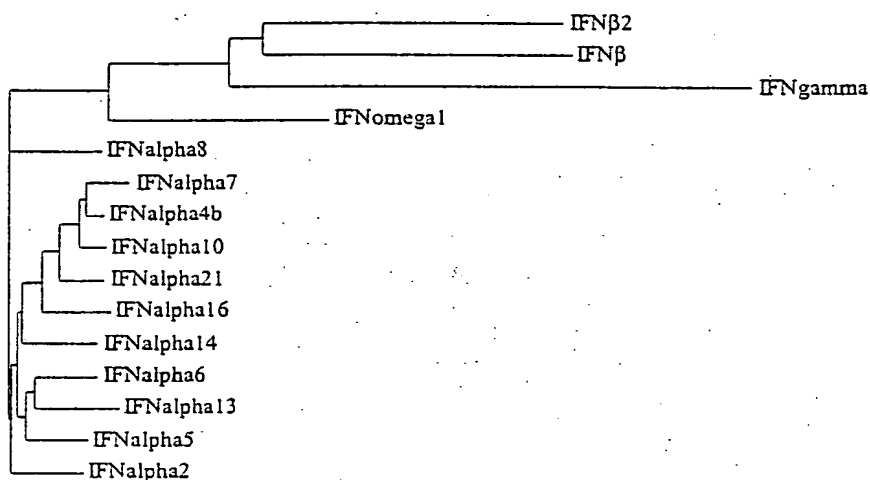
Figure 4

MAIL DATE 23-02-2002
O I P E J C 6 1
PATENT & TRADEMARK OFFICE

09004050.1125

NOV 25 2002
O I P E J C 6 1
PATENT & TRADEMARK OFFICE

Protein Level Comparison



Nucleotide Level Comparison

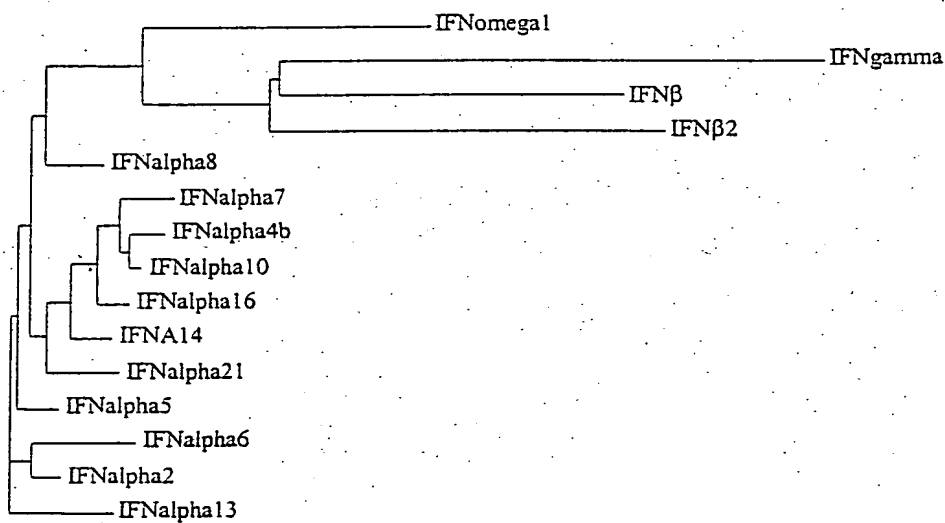


Figure 5

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DATE 25 2002
PATENT & TRADEMARK

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NOV 25 2002

TRADEMARK OFFICE

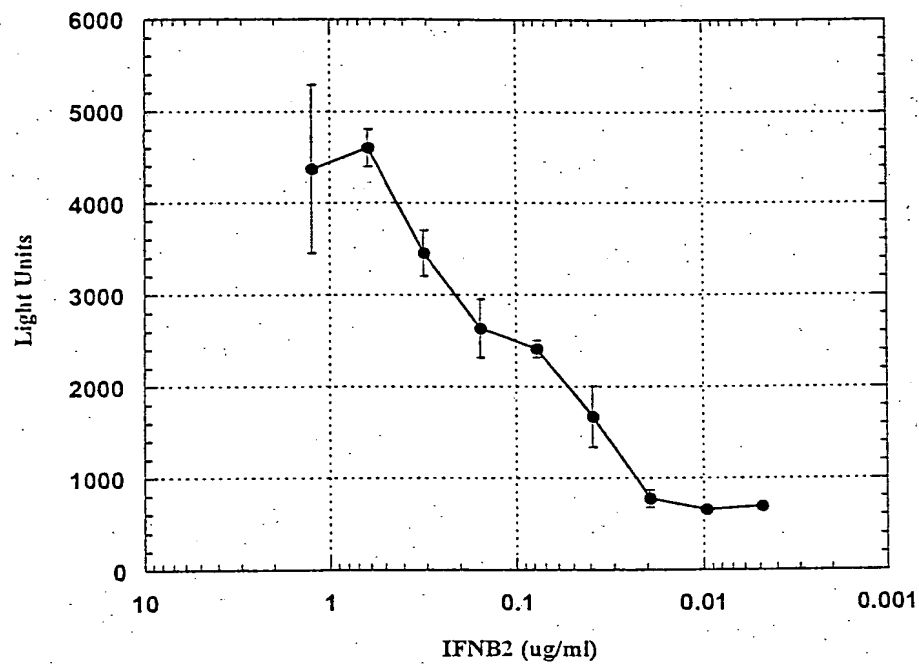


Figure 6



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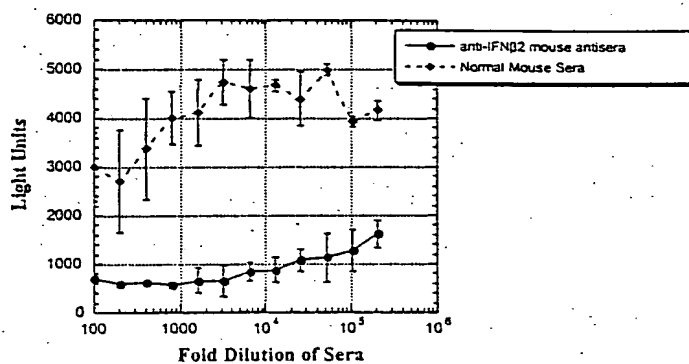
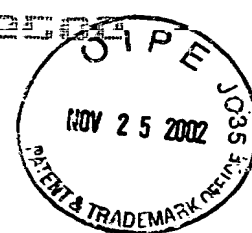


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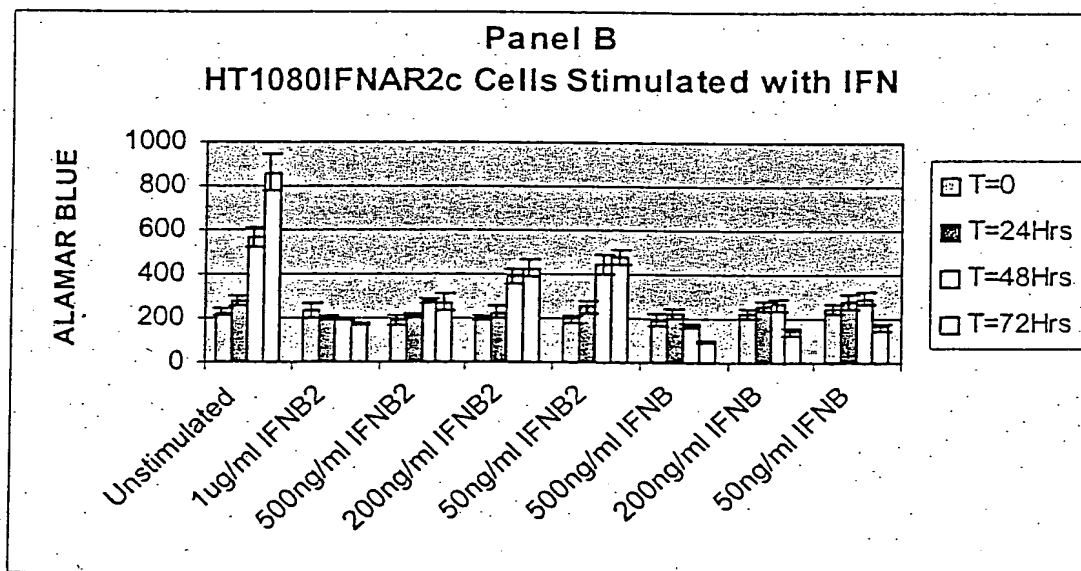
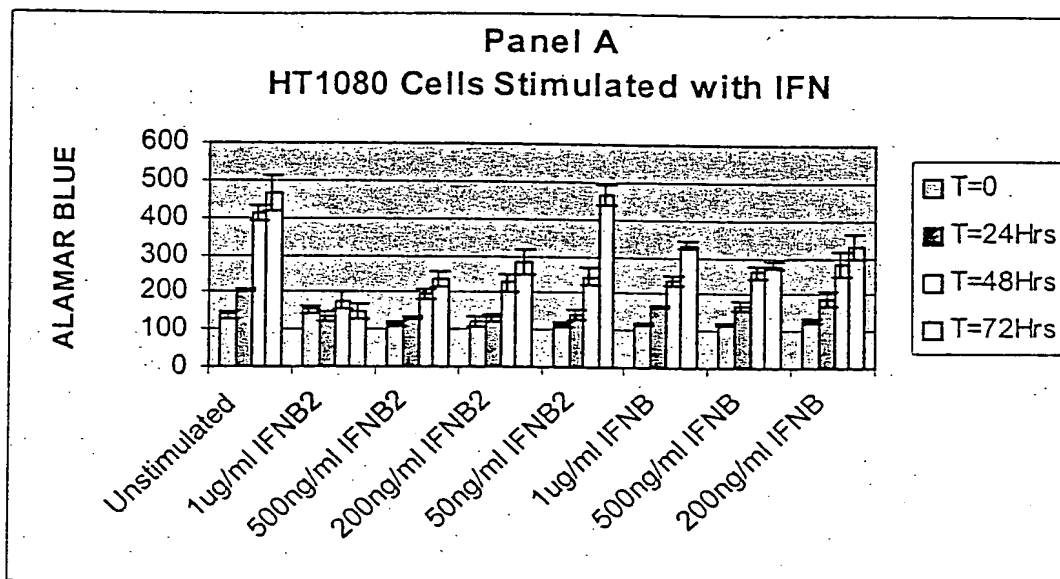
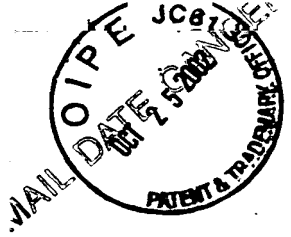


Figure 8



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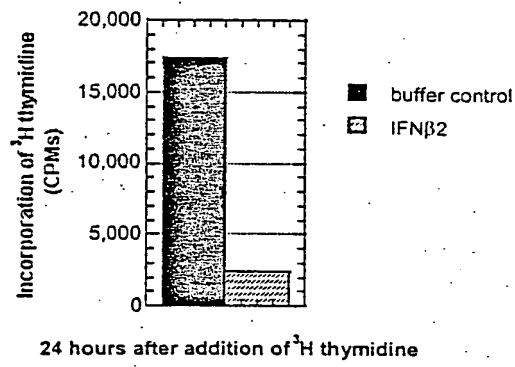
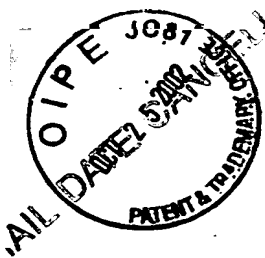


Figure 9



09001050 . 112502

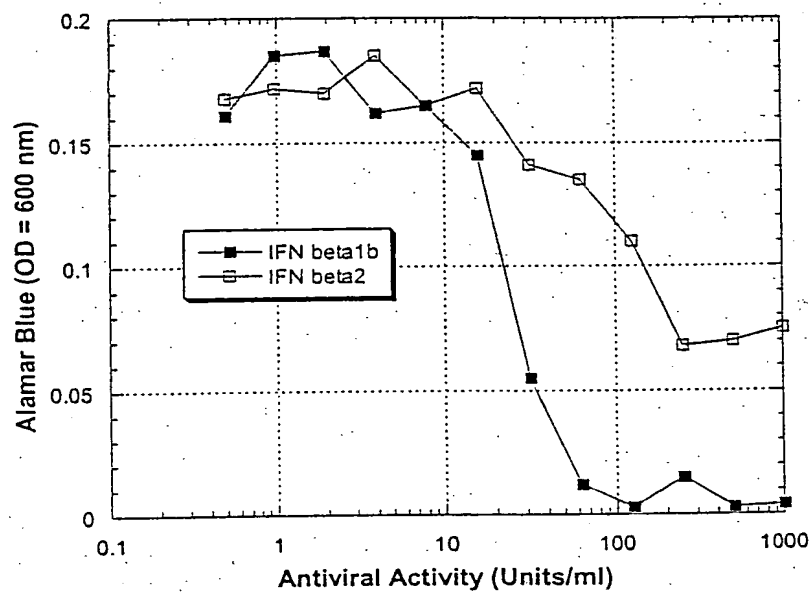
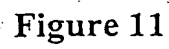
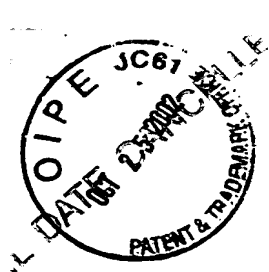


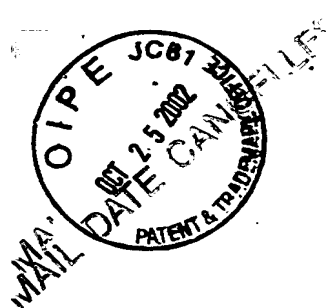
Figure 10





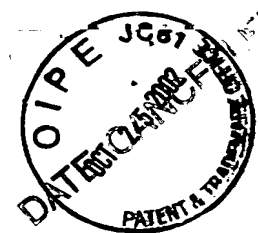
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181 acacaatagg gaaatggaga aaatgtactc tgaacaccat gaaaagggaa cctgaaaatc
241 taatgtgtaa acttggagaa atgacattag aaaacgaaag ctacaaaaga gaacactctt
301 caaaataatc tgagatgcat gaaaggcaaa cattcactag agctggaatt tccttaagtc
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Figure 12



361 atgatt atcaagcact tctttggaac
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481 ccagcaaaga caagtgaatc aagaaagttt aaaactcttg aataagttgc aaaccttgtc
541 aattcagcag tgtctaccac acaggaaaaa ctttctgctt cctcagaagt ctttgagtcc
601 ttaactgtac caaaaaggac acactctggc cattcttcat gagatgct

Figure 13



09881050.112507



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Figure 14

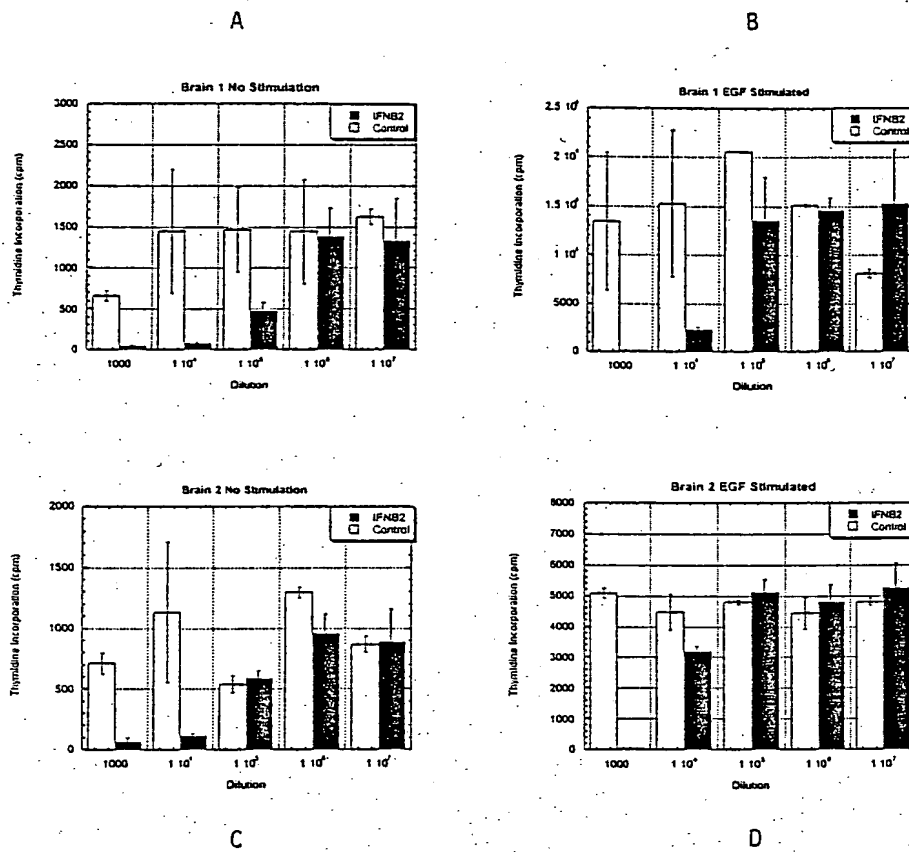
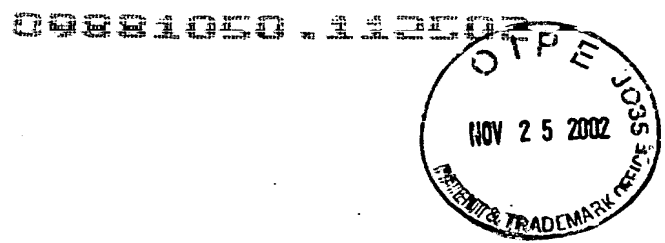
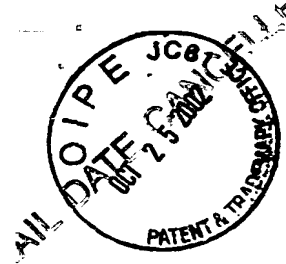


Figure 15